* *
Reviewer Comments:

<110> Jackson, Aimee L.
       Bartz, Steven R.
       Burchard, Julja,
       Linsley, Peter S.
       Ge, Wei
       Cavet, Guy L.

<120> METHOD OF DESIGNING siRANS FOR GENE
   SILENCING

Please change "siRANS" in the above <120> response to "siRNAS"

<210>
335 <211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA sequence corresponding to siRNA sense strand
       in construction of siRNA library

Please move the above "335" up to the <210> line.
* *

Application No: 10577696   Version No: 1.0

Input Set:

Output Set:

Started: 2007-12-14 12:08:13.705
Finished: 2007-12-14 12:08:19.590
Elapsed: 0 hr(s) 0 min(s) 5 sec(s) 885 ms
Total Warnings: 990
Total Errors: 2
No. of SeqIDs Defined: 990
Actual SeqID Count: 990

Error code   Error Description

W 213   Artificial or Unknown found in <213> in SEQ ID (1)

W 213   Artificial or Unknown found in <213> in SEQ ID (2)

W 213   Artificial or Unknown found in <213> in SEQ ID (3)

W 213   Artificial or Unknown found in <213> in SEQ ID (4)

W 213   Artificial or Unknown found in <213> in SEQ ID (5)

W 213   Artificial or Unknown found in <213> in SEQ ID (6)

W 213   Artificial or Unknown found in <213> in SEQ ID (7)

W 213   Artificial or Unknown found in <213> in SEQ ID (8)

W 213   Artificial or Unknown found in <213> in SEQ ID (9)

W 213   Artificial or Unknown found in <213> in SEQ ID (10)

W 213   Artificial or Unknown found in <213> in SEQ ID (11)

W 213   Artificial or Unknown found in <213> in SEQ ID (12)

W 213   Artificial or Unknown found in <213> in SEQ ID (13)

W 213   Artificial or Unknown found in <213> in SEQ ID (14)

W 213   Artificial or Unknown found in <213> in SEQ ID (15)

W 213   Artificial or Unknown found in <213> in SEQ ID (16)

W 213   Artificial or Unknown found in <213> in SEQ ID (17)

W 213   Artificial or Unknown found in <213> in SEQ ID (18)

W 213   Artificial or Unknown found in <213> in SEQ ID (19)

W 213   Artificial or Unknown found in <213> in SEQ ID (20)

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              Input Set:


             Output Set:



               Started:   2007-12-14 12:08:13.705
              Finished:   2007-12-14 12:08:19.590
               Elapsed:   0 hr(s) 0 min(s) 5 sec(s) 885 ms
        Total Warnings:   990
          Total Errors:   2
   No. of SeqIDs Defined:  990
      Actual SeqID Count:  990
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Error code	Error Description
	This error has occured more than 20 times, will not be displayed
E 201	Mandatory field data missing in <210> Expected SEQID 335
E 212	Invalid Sequence ID Number; Expected 335 as next SeqID but skipped

<210> 1
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<220>
<223> an example of siRNA

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<210> 2
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<220>
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      in construction of siRNA library

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<210> 3
<211> 19

<212> DNA
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<220>
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      in construction of siRNA library

<400>   3
actcttactg ctctccagt   19

<210>   4
<211> 19
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<213> Artificial Sequence

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      in construction of siRNA library

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cttaacacgg atgctggtg   19

<210> 5
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      in construction of siRNA library

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ggagagcttt ctaggacct   19

<210> 6
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      in construction of siRNA library

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agtcatcccg cagagccgc   19

<210> 7
<211> 19
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      in construction of siRNA library

<400> 7
atcgtagtgc ttgtactta   19

<210> 8
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<220>
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<210> 9
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<210> 10
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<210> 11
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<220>
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<400> 11
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<210> 12
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<220>
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<210> 14
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<210> 16
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ggcctctata cccctcaaa   19

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<210> 19
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<400> 19
cagagacaga atctacact   19

<210> 20
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<400> 20
caacagaagg ttgtcttgt   19

<210> 21
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<400> 21
ttgtgtgtgg gaccgtaat   19

<210> 22
<211> 19

<212> DNA
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<220>
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<210> 23
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<400> 23
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<210> 24
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<220>
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<210> 25
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<220>
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<400> 25
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<210> 26
<211> 19
<212> DNA
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<220>
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<400> 26
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<210> 27
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<210> 28
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<210> 29
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<220>
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<210> 30
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<220>
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<400> 30
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<210> 31
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<220>
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<400> 31
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<210> 32
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<220>
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<400> 32
cattactcgg ggggccatc   19

<210> 33
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<220>
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<400> 33
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<210> 34
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<220>
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<210> 35
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<220>
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<400> 35
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<210> 36
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<220>
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<400> 36
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<210> 37
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
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<400> 37
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<210> 38
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<220>
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<400> 38
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<210> 39
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<400> 39
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<210> 40
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<210> 41
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<212> DNA
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<400> 41
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<210> 43
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<400> 43
taattcacag ggacctaaa   19

<210> 44
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<400> 44
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<210> 45
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<220>
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<400> 45
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<210> 46
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<210> 47
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<220>
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<210> 48
<211> 19
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<220>
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<210> 49
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<220>
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<210> 51
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<210> 52
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<210> 53
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<210> 54
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<220>
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<400> 54
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<210> 55
<211> 19
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<400> 55
gacctgtgcc ttttagaga   19

<210> 56
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<210> 57
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<210> 58
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<210> 59
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<400> 59
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<210> 60
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<210> 62
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<220>
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<210> 68
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<210> 69
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<210> 70
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<220>
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<400> 70
ctctaccact gaagagttg   19

<210> 71
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<220>
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